

<213> Artificial Sequence

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SEQUENCE LISTING

RECEIVED <110> The University of Queensland MAY 0 3 2002 <120> Novel omega donotoxin peptides <130> 2338740/MJC **TECH CENTER 1600/2900** <140> US 09/679,490 <141> 1999-04-16 PCT/AU99/00288 <150> <151> 1999-04-16 <160> 44 <170> PatentIn version 1.0 <210> 1 <211> <212> PRT <213> conus catus <400> 1 Ser Gly Thr Val Gly Arg <210> 2 <211> 6 <212> PRT <213> conus catus <400> 2 Ser Lys Leu Met Tyr Asp <210> 3 <211> 6 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: A modified version of the second loop of CVID <400> 3 Ser Arg Leu Met Tyr Asp <210> <211> 6 <212> PRT

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<220>
<223> Description of Artificial Sequence: A modified version of the second
      loop of CVID
<400> 4
Asp Arg Leu Met Tyr Asp
<210> 5
<211> 27
<212> PRT
<213> conus catus
<400> 5
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
               5
                            10
Ser Gly Ser Cys Ser Gly Thr Wal Gly Arg Cys
           20
<210> 6
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A modified form of CVID
<400> 6
Cys Lys Ser Lys Gly Ala Lys Cys Ser Arg Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
           20
<210> 7
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A modified form of CVID
<400> 7
Cys Lys Ser Lys Gly Ala Lys Cys Asp Arg Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg/ Cys
           20
<210> 8
<211> 25
<212> PRT
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<213> conus magus
<400> 8
Cys Lys Gly Lys Gly Ala Lys Cys Ser Arg Leu Met Tyr Asp Cys Cys
Thr Gly Ser Cys Arg Ser Gly Lys Cys
           20
<210> 9
<211> 26
<212> PRT
<213> conus magus
<400> 9
Cys Lys Gly Lys Gly Ala Pro Cys Arg Lys Thr Met Tyr Asp Cys Cys
Ser Gly Ser Cys Gly Arg Arg Gly Lys Cys
           20
<210> 10
<211> 27
<212> PRT
<213> conus geographus
<220>
<221> misc_feature
<223> Pro at positions 4, 10 and 21 is 4-Hyp
<400> 10
Cys Lys Ser Pro Gly Ser Ser Cys Ser Pro Thr Ser Tyr Asn Cys Cys
Arg Ser Cys Asn Pro Tyr Thr Lys Arg Cys Tyr
<210> 11
<211> 18
<212> DNA
<213> conus catus
<400> 11
                                                                     18
agcggcaccg taggtaga
<210> 12
<211> 382
<212> DNA
<213> conus catus
<220>
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<221> CDS

<222> (10)..(228) <400> 12 atcatcaaa atg aaa ctg acg tgt gtg gtg atc gtc gcc gtg ctg ctc ctg Met Lys Leu Thr Cys Val Val Ile Val Ala Val Leu/Leu Leu acg gcc tgt caa ctc atc aca gct aat gac tcc aga ggt acg cag aag 99 Thr Ala Cys Gln Leu Ile Thr Ala Asn Asp Ser Arg Gly/Thr Gln Lys 20 25 cat cgt gcc ctg agg tcg gac acc aaa ctc tcc atg/tcg act cgc tgc 147 His Arg Ala Leu Arg Ser Asp Thr Lys Leu Ser Mot Ser Thr Arg Cys aag agt aaa gga gca aaa tgt tca aag ctt atg tat gac tgc tgc agc 195 Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys Ser 55 ggt tot tgc agc ggc acc gta ggt aga t/gt ggc tgatccggcg cttgatctcc 248 Gly Ser Cys Ser Gly Thr Val Gly Arg/Cys Gly 70 cccttctgtg ctctatcctt ttctgcctga/gtcctcctta cctgagagtg gtcatgaacc 308 actcatcacc tacccctgg aggtctcaaa gaactacttg aaataaagcc gcttgcaaaa 368 aaaaaaaaa aaaa 382 <210> 13 <211> 73 <212> PRT <213> conus catus <400> 13 Met Lys Leu Thr/Cys Val Val Ile Val Ala Val Leu Leu Leu Thr Ala Cys Gln Leu/Ile Thr Ala Asn Asp Ser Arg Gly Thr Gln Lys His Arg 25 Ala Leu Arg Ser Asp Thr Lys Leu Ser Met Ser Thr Arg Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys Ser Gly Ser 55 60 Cýs Ser Gly Thr Val Gly Arg Cys Gly

<210> 14 <211> 27 <212> PRT

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<213> conus catus
<400> 14
Cys Arg Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp/Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210> 15
<211> 27
<212> PRT
<213> conus catus
<400> 15
Cys Lys Ser Lys Gly Ala Arg Cys Ser Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
            20
<210> 16
<211> 27
<212> PRT
<213> conus catus
<400> 16
Cys Lys Ser Lys Gly Ala/Gln Cys Ser Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
           20
<210> 17
<211> 27
<212> PRT
<213> conus catús
<400> 17
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Ala Val Gly Arg Cys
           20
<210>
<211>
<212>
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<213>
<220>
      Description of Artificial Sequence: A derivative of CVID
<40,0>
     18
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Cys Lys Ser Lys Gly Ala Lys Cys Asp Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210> 19
<211>
      27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A derivative of CVID
<400> 19
Cys Lys Tyr Lys Gly Ala Lys Cys Ser Arg Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
           20
<210> 20
<211> 27
<212> PRT
<213> Artificial Segmence
<220>
<223> Description/of Artificial Sequence: A derivative of CVID
<400> 20
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Ala Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210>
<211>
<212>
<213>
      Ártificial Sequence
<220>
      Description of Artificial Sequence: A derivative of CVID
<223%
<400> 21
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
Thr Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210> 22
<211> 27
<212> PRT
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: A derivative of CVID
<220>
<221> misc_feature
<223> Xaa at position 5 is D-alanine
<400> 22
Cys Lys Ser Lys Xaa Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
                                    10
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cy
<210> 23
<211>
      28
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial/Sequence: A derivative of CVID
<400> 23
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys Tyr
            20
<210> 24
<211>
      27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A derivative of CVID
<400> 24
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser/Cys Ser Gly Thr Val Gly Arg Cys
<210>
       25
<211>
       2,8
<212>
       PRT
      Artificial Sequence
<213>
      Description of Artificial Sequence: A derivative of CVID
<400> 25
Tyr Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys
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Cys Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210> 26
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A dérivative of CVID
<220>
<221> misc feature
<223> Cysteine at position 1 is acylated
<400> 26
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
                                   10
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
            20
                                25
<210> 27
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A derivative of CVID
<220>
<221> misc_feature
<223> Leu at position 12 is L-norleucine
<400> 27
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Leu Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
           20
<210>
      /28
<211>
      27
<212>
      PRT
<213
      Artificial Sequence
<220'>
<22β> Description of Artificial Sequence: A derivative of CVID
<220>
<221> misc feature
<223> Leu at position 12 is L-norleucine
k400> 28
Cys Lys Ser Lys Gly Ala Lys Cys Ser Arg Leu Leu Tyr Asp Cys Cys
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Ser Gly Ser\Cys Ser Gly Thr Val Gly Arg Cys
<210> 29
<211>
      27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A derivative of CVID
<220>
<221> misc_feature
<223> Leu at position 12 is L-norleucine
<400> 29
Cys Lys Tyr Lys Gly Ala Lys Cys Ser Arg Leu Leu Tyr Asp Cys Cys
               5
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
            20
<210> 30
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A derivative of CVID
<220>
<221> misc_feature
<223> Xaa at position 12 is L-O-methyl homoserine
<400> 30
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Xaa Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
<210> 31
<211> 27
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A derivative of CVID
<220>
<221> misc_feature
<223> Methionine residue at position 12 is oxidised to its sulfoxide
<400> 31
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Xaa Yyr Asp Cys Cys
                                   10
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Ser Gly Ser Cys Ser Gly Thr\Val Gly Arg Cys
           20
                               25
<210> 32
<211>
      27
<212>
      PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A derivative of CVID
<400> 32
Cys Lys Ser Lys Gly Ala Lys Cys Ser Lys Leu Met Tyr Asp Cys Cys
Ser Gly Ser Cys Ser Gly Thr Val Gly Arg Cys
           20
<210> 33
<211>
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A modified version of the second loop
<400> 33
Asp Lys Leu Met Tyr Asp
<210> 34
<211>
<212>
      PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A modified version of the second loop
       of CVID
<400> 34
Ser Lys Leu Ala Tyr Asp
<210> 35
<211>
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A modified version of the second loop
       of CVID
<220>
<221> misc_feature
<223> Leu at position 4 is L-norleucine
<400> 35
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Ser Lys Leu Leu Tyr Asp
<210> 36
<211> 6
<212> PRT
<213> Artificial $equence
<220>
<223> Description of Artificial Sequence: A modified version of the second loop
      of CVID
<220>
<221> misc feature
<223> Leu at position 4 \is L-norleucine
<400> 36
Ser Arg Leu Leu Tyr Asp
<210> 37
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A modified version of the second loop
      of CVID
<220>
<221> misc_feature
<223> Xaa at position 4 is L-O-methyl homoserine
<400> 37
Ser Lys Leu Xaa Tyr Asp
<210> 38
<211> 6
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: A modified version of the second loop
       of CVID
<220>
<221> misc_feature
<223> Xaa at position 4 is L-O-methyl serine
<400> 38
Ser Lys Leu Xaa Tyr Asp
```

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<210>
<211>
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer
<220>
<221> misc_feature
<223> Xaa may be any other amino acid and up to one Xaa may be a
      deletion
<400> 39
Cys Xaa Xaa Xaa Gly Xaa Xaa Cys Xaa Lys Leu Xaa Tyr Xaa Cys Cys
Xaa Ser Cys Ser Gly Xaa Val Gly Arg Cys
<210> 40
<211>
      28
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: primer
<400> 40
aactggaaga attcgcggcc gcaggaat
                                                                     28
<210> 41
<211>
      23
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer
<400> 41
atcatcaaaa tgaaactgac gtc
                                                                     23
<210> 42
<211>
      28
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer
<400> 42
                                                                     28
aactggaaga attcgcggcc gcaggaat
```

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<210>
<211>
      27
<212>
      DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer
<400> 43
atcaaaatga aactgadgtg tgtggtg
                                                                     27
<210>
<211>
      26
<212>
<213>
      Artificial Sequence
<220>
      Description of Artificial Sequence: primer
<400> 44
gcgttttgat cagccacatc taccta
                                                                     26
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